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		_	,	Aud				
1: AA212542. mu44c08.r1 Soares[gi:1811196] Links								
LOCUS DEFINITION		.rl Soare	s_thymus_2N			inear EST 19 cDNA clone IMA	-FEB-1997 .GE:642254	
ACCESSION	5', mRNA sequence. AA212542 AA212542.1 GI:1811196							
VERSION KEYWORDS	EST.							
SOURCE Mus musculus (house mouse) ORGANISM Mus musculus Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.								
								REFERENCE AUTHORS
	Theisin	g,B., Wyl:	ie,T., Lenn	on, G., S	oares,B.,	Wilson, R. and	· •	
TITLE	Waterston,R. The WashU-HHMI Mouse EST Project							
JOURNAL COMMENT	Unpublished (1996) Contact: Marra M/Mouse EST Project							
	WashU-HHMI Mouse EST Project Washington University School of MedicineP 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108 Tel: 314 286 1800 Fax: 314 286 1810 Email: mouseest@watson.wustl.edu This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information. MGI:394246							
	Seq primer: -28m13 rev2 ET from Amersham High quality sequence stop: 461.							
FEATURES	High qu		dence stop: n/Qualifier					
source		1470 /organi:	sm="Mus mus	culus"				
/strain="C57BL/6J" /db_xref-" <u>taxon:10090</u> " /clone="IMAGE:642254"								
/clone_lib="Soares_thymus_2NbMT" /sex="male"								
/tissue_type="Thymus" /dev stage="4 weeks"								
<pre>/lab_host="DH10B" /note="Vector: pT7T3D-Pac (Pharmacia) with a modified</pre>							fied	
polylinker; Site_1: Not I; Site_2: Eco RI; 1st strand cDNA was primed with a Not I - oligo(dT) primer [5' TGTTACCAATCTGAAGTGGGAGCGGCCGCGTTTTTTTTTT								
							the Not I	
							A	
		rounds o		ation, a	nd was cor	nstructed by B		
BASE COUNT	130							

ORIGIN

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1 gtgatggtgc tatttattat tgcctttgag ccatctaccc tgattatcct atacacattt 61 cttgtgctgt cataaaaata agatacagta actgcatgct tgctggtggg tacccagttc 121 ttctttgtgt tcgggtcaat ctggaagaca tgagctcgag tgctgaagat aggttgctcc 181 cccattttgc ccaatgaaaa cttgctctga agtttctgct cttatgctac agaaataact 241 tccaaggaat tttctccgtc tgttattttg cagttgcttt tccacccccg agatcttttc 301 cggagcaact caacaaggca gacaattcag ttagttcttg tcggtaaaat gatttcactc 361 gcgctgacac caaaatgttc aaacatttgg aatttgaaaa ggttgcgcgg agaatatagc 421 acttaactat tccacaaaat taagcgtacg tgtagaaaaa acgcaaaaagc
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